Sequence Listing

<110> Sidhu, Sachdev S.
Weiss, Gregory A.
Wells, James A.

<120> TRANSFORMATION EFFICIENCY IN PHAGE DISPLAY THROUGH MODIFICATION OF A COAT PROTEIN

<130> 146392004400

<140> US 09/380,447
<141> 1999-09-01

<150> US 60/134,870
<151> 1999-05-19

<150> US 60/133,296
<151> 1999-05-10

<150> US 60/103,514
<151> 1998-10-08

<150> US 60/094,291
<151> 1998-07-27

<150> PCT/US99/16596
<151> 1999-07-22

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Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa

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Asp Asp Gly Glu Ala
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Val Ile Val Gly Ala Thr Ile Gly Ile Lys Leu Phe Lys Lys Phe
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Thr Ser Lys Ala Ser
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Thr Ser Lys Ala Ser
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Asp Asp Gly Glu Ala
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<400> 68
gaacttttct ttctcgcggg gactgtgcat ctt 33

<210> 69
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<400> 69
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gcaaaagcgg cctataactc ccttgaggat attgct 36

<210> 75
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<400> 75
gcaaaagcgg cctataacgc tcttgaggat tcagctacta actatatc 48

<210> 76
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<220>
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<400> 76

cccgcaaaag cggcctatga gtcccttgag gattcagcta ctaactatat 50

cggttatgcg 60

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gcaaaagcgg cctataactc ccttgaggat tcagctacta actatatc 48

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Gln Ser Gly Gly Gly Ser Gly Ser Ser Ser
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<210> 79
<211> 5
<212> PRT
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<220>
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<400> 79
Gly Gly Arg Pro Val
1 5

<210> 80
<211> 34
<212> DNA
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<220>
<223> linker oligonucleotide

<400> 80
cagagcggtg gaggatccgg gagctccaga gggt 34

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<212> DNA
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cagagcggtg gaggatccgg gagctccagc gccgagggt 39

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<220>
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<400> 82
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<210> 83
<211> 60
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<220>
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<400> 83
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ccatcaccat 60

<210> 84
<211> 60
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<400> 84
gctgtcggta ttatttacat gctcctcgtg gaggcgtcgc cctgggctgc 50

taaggcgcca 60

<210> 85
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<400> 86
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<210> 87
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<400> 87
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tgcg 54

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tcaccatgcg 60

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ccatcaccat gcg 63

<210> 95
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ccatcaccat caccatgcg 69

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tggtggccat caccatcacc atgcg 75

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tggcggtggt ggccatcacc atcaccatgc g 81

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tgttgat 57

<210> 102
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ttggatttgg gctgtcggt 69

<210> 103
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gttttttctg ttgatgttga tnnsnnsnns nnsnnsnnsn nsnnsnnsnn 50

sgcggctgat gcattccca 69

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tgctaaggcg ccagacgatg gt 72

<210> 105
<211> 69
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agcgctcagc tgagcaactt cnnsnnsnns nnsnnsnnsn nsnnsnnsnn 50

sgcggctgat gcattccca 69

<210> 106
<211> 81

<212> DNA
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<220>
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<400> 106
gatggtgaag ctgcggctvv cvvcvvcvvc vvcvvcvvcv vcvvcvvcvv 50

cvvcvvcvvc gatgcattcc caactatacc a 81

<210> 107
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tnwtnkgnyt nkgnytnwcn ktnwtnwtga gactgctagc gctcag 96

<210> 108
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<400> 108
caccatcacc atcaccatgc g 21

<210> 109

<211> 30
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<400> 109
gcctgggagg agaacatcga cagcgccccc 30

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<220>
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<400> 110
Ala Trp Glu Glu Asn Ile Asp Ser Ala Pro
1 5 10

<210> 111
<211> 30
<212> DNA
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<220>
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<400> 111
cagtacggga cgccggacac cgacaccgac 30

<210> 112
<211> 10
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Gln Tyr Gly Thr Pro Asp Thr Asp Thr Asp
1 5 10

<210> 113
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acggggtggt tggaggggcc cgacaccccc 30

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<400> 114
Thr Gly Trp Leu Glu Gly Pro Asp Thr Pro
1 5 10

<210> 115
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<220>
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<400> 115
ctcatgggcc ccggcgcgga cggc 24

<210> 116
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<220>
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Leu Met Gly Pro Gly Ala Asp Gly
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<210> 117
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<400> 117
cacgactcgg tcccgagcaa cggc 24

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<400> 118
His Asp Ser Val Pro Ser Asn Gly
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<210> 119
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<220>
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gctgagcaac ttcgctgcta aggcgccaga cgatggtgaa gctgcggctc 100

accatcacca tcaccatgcg 120

<210> 120
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Ala Gln Leu Ser Asn Phe Ala Ala Lys Ala Pro Asp Asp Gly Glu
20 25 30

Ala Ala Ala His His His His His His Ala
35 40

<210> 121
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<212> PRT
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<220>
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<400> 121
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1 5 10 15

Ser Ala Gln Leu Ser Asn Phe Ala Ala Lys Ala Pro Asp Asp Gly
20 25 30

Glu Ala Ala Ala His His His His His His Ala
35 40

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<211> 51
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<400> 122
gctgcggctg atgcatctgg tagcgtctag agccaccatc accatcacca 50

t 51

<210> 123
<211> 54
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<220>
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<400> 123
Met Ser Lys Ser Thr Phe Lys Lys Phe Leu Lys Val Phe Val Phe
1 5 10 15

Ser Val Asp Val Asp Asn Asn Trp Ile Trp Ala Val Gly Ile Ile
20 25 30

Glu Thr Ala Ser Ala Gln Leu Ser Asn Phe Ala Ala Lys Ala Pro
35 40 45

Asp Asp Gly Glu Ala Ala Ala Asp Ala
50

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<211> 150
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<220>
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tgatgttgat aataattgga tttgggctgt cggtattatt tacatgctcc 100

tcgtggaggc gtcgccctgg gctgctaagg cgccagacga tggtgaagct 150

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t 51

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cggcaccggc 60

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Ala Arg Gly Thr Gly
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cggcagccac 60

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Ala Arg Ser Gly Pro
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Arg Gly Ser Asn Gly Ser Asp Ser Ser Ser
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tattgtt 57

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1 5 10 15

Val Gly Ile Val

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ttatgtt 57

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1 5 10 15

Tyr Gly Tyr Val

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tcttgtt 57

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Leu Phe Leu Val

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ttttgat 57

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Asn Ser Phe Asp

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tgttaat 57

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Val Gly Ala Thr Ile Gly Ile Lys Leu Phe Lys Lys Phe Thr Ser Lys
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Val Gly Ala Thr Ile Gly Ile Lys Leu Phe Lys Lys Phe Thr Ser Lys
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Ala Ser
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            20                  25                  30
Val Gly Ala Thr Ile Gly Ile Lys Leu Phe Lys Lys Phe Thr Ser Lys
        35                  40                  45
Ala Ser
    50
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